EMERGING AND RE-EMERGING INFECTIOUS DISEASES

Despite remarkable advances in medical research and treatments during the 20th century, infectious diseases remain among the leading causes of death worldwide for three reasons: (1) emergence of new infectious diseases, (2) re-emergence of old infectious diseases, and (3) persistence of intractable infectious diseases. Emerging diseases include outbreaks of previously unknown diseases or known diseases whose incidence in humans has significantly increased in the past two decades. Re-emerging diseases are known diseases that have reappeared after a significant decline in incidence. Within the past two decades, innovative research and improved diagnostic and detection methods have revealed a number of previously unknown human pathogens. (For a list of emerging and re-emerging infectious diseases and pathogens, see the table below or www.niaid.nih.gov/dmid/eid/erd.htm.) Largely as a result of better detection methods, evidence also is accumulating that infective agents play a role in diseases previously thought to be chronic and noncommunicable. For example, during the past decade, chronic gastric ulcers, which formerly were thought to be caused by stress or diet, were found to be the result of infection by the bacterium Helicobacter pylori.

New infectious diseases continue to evolve and "emerge." Changes in human demographics, behavior, land use, and other factors are contributing to new disease emergence by changing transmission dynamics to bring people into closer and more frequent contact with pathogens. This situation may involve exposure to animal or arthropod carriers of disease. Increasing trade in exotic animals for pets and as food sources has increased opportunities for pathogens to jump from animal reservoirs to humans. For example, close contact with exotic rodents imported to the United States as pets led to the recent outbreak of monkeypox in this country, and use of exotic civet cats for meat in China

was found to be one potential route by which the severe acute respiratory syndrome (SARS) coronavirus made the transition from animal to human hosts.

In addition to the continual discovery of new human pathogens, old infectious disease enemies are "re-emerging." Natural genetic variations, recombinations, and adaptations allow new strains of known pathogens to appear. The immune system has not been previously exposed to these new strains and therefore is not primed to recognize them (e.g., influenza). Furthermore, human behavior plays an important role in disease re-emergence. Increased and sometimes imprudent use of antimicrobial drugs and pesticides has led to the development of resistant pathogens, allowing many diseases that were formerly treatable with drugs to make a comeback (e.g., tuberculosis, malaria, hospitalacquired and food-borne infections). Recently, decreased compliance with vaccination policy also has led to re-emergence of diseases such as measles and pertussis, which were previously under control. The use of deadly pathogens such as smallpox or anthrax as agents of bioterrorism is an increasingly acknowledged threat to the civilian population. Moreover, many important infectious diseases have never been adequately controlled on either the national or international level. Infectious diseases that have posed ongoing health problems in developing countries such as food- and water-borne infections, dengue, and West Nile virus, are re-emerging in the United States.

NIAID has developed a strategy to address the threat of emerging and re-emerging infectious diseases through targeted research and training. That strategy, which was initially outlined in the Institute's 1996 document, *The NIAID Research Agenda for Emerging Infectious Diseases (www. niaid.nih.gov/publications/execsum/bookcover.htm)*, was updated in the 2000 NIAID strategic plan, *NIAID: Planning for the 21st Century (www.niaid. nih.gov/strategicplan/pdf/splan.pdf)*. In May 2001, NIAID released *NIAID Global Health Research*

Plan for HIV/AIDS, Malaria, and Tuberculosis (www.niaid.nih.gov/publications/globalhealth/global.pdf). This document outlines the Institute's plans for the next decade for diagnosing, treating, and preventing these three infections and also lays out a plan for enhancing in-country research capacity.

To an unprecedented extent, issues related to global health and infectious diseases are on the agendas of world leaders, health policymakers, and philanthropies. This attention has been focused both on scientific challenges such as vaccine development, and on the deleterious effects of infectious diseases on economic development and political stability.

To enhance the capacity to deal with the challenges posed by emerging diseases, NIAID opened a new biosafety level 3 (BSL-3) laboratory in Hamilton, Montana, in April 2002, and anticipates completion of a new BSL-3 laboratory in Rockville, Maryland. NIAID has also broken ground on a new BSL-3 laboratory on the main NIH campus in Bethesda, Maryland. These facilities will enable the Institute to conduct animal studies and laboratory research on infectious agents such as drug-resistant *Mycobacterium tuberculosis* (*M.tb*), *Borrelia*, *Yersinia*, the influenza virus, West Nile, dengue virus, and other important pathogens.

Likewise, in fiscal year (FY) 2003 NIAID established several new extramural programs to address needs for biodefense and emerging infectious disease research:

National Biocontainment Laboratories (NBLs) and Regional Biocontainment Laboratories (RBLs). NIAID has provided funding for the construction of several new biocontainment facilities for biodefense and emerging infectious disease research: two NBLs that will provide BSL-4 research facilities (construction is expected to take up to 5 years); and nine RBLs that will provide BSL-3 research facilities (construction of

- these facilities is expected to take from 2 to 3 years).
- Regional Centers of Excellence (RCEs) for Biodefense and Emerging Infectious Diseases Research. Eight RCE grants were funded in September 2003, for centers located throughout the country. These RCEs are consortia of institutions usually within a region. They are funded to develop and maintain strong infrastructure and multifaceted research and development activities that will provide the scientific information and translational research capacity to make the next generation of therapeutics, vaccines, and diagnostics against the NIAID Category A, B, and C agents. Two planning grants for RCEs were funded to enable institutions to develop consortia for the study of these pathogens and to support research program initiation and resource acquisition.
- Infections Research Resources Program supports the acquisition, authentication, storage, and distribution to the scientific community of state-of-the-art research and reference reagents related to biodefense and emerging infectious diseases. Included are the capability to validate, expand, and produce biological agents, including cell lines, clones, proteins, monoclonal and polyclonal antibodies, and diagnostic tools.
- Ocontracts funded under the *In Vitro* and Animal Models for Emerging Infectious Diseases and Biodefense program provide a range of resources for preclinical testing of new therapies and vaccines, including nonhuman primate models. Under these new contracts, task orders for small animal and nonhuman primate models for anthrax are being developed and validated for Food and Drug Administration (FDA) licensure of vaccines and therapeutics against inhalational anthrax. Included in this activity are safety,

toxicology, and pharmaceutical testing in small and large animals, including the capability for conducting challenge studies. Future task order awards will develop, validate, and use models for other Category A, B, and C pathogens.

Integrated Research Network expands the Institute's capacity to conduct clinical research studies of food- and water-borne enteric pathogens. Through one of the Microbiology Research Units with a botulism research center, NIAID is determining the pharmacokinetics of botulinum toxins and antitoxins in several animal models. These data are helpful for the development, formulation, and use of equine-based antitoxins and will serve as a basis for next-generation antitoxins.

Basic and clinical research is critical to the development of a national strategy to confront these microbial challenges. Such research increases our collective understanding of everchanging microbial populations and permits this new knowledge to be transformed into better diagnostics, vaccines, and therapies. Basic research and research training also are the foundation for surveillance and response activities.

During 2004, NIAID supported research initiatives on biodefense as well as on emerging and re-emerging infectious diseases in multiple areas, including SARS, influenza, TB, and other infectious diseases and deadly pathogens.

List of NIAID Emerging and Re-emerging Diseases 2004

Group I—Pathogens Newly Recognized in the Past Two Decades

Acanthamebiasis

Australian bat Lyssavirus

Babesia, atypical

Bartonella henselae

Coronaviruses/Severe Acute Respiratory Syndrome (SARS)

Fhrlichiosis

Encephalitozoon cuniculi

Encephalitozoon hellem

Enterocytozoon bieneusi

Helicobacter pylori

Hendra or equine morbilli virus

Hepatitis C

Hepatitis E

Human herpesvirus 8

Human herpesvirus 6

Lyme borreliosis

Microsporidia

Parvovirus B19

Group II—Re-emerging Pathogens

Coccidioides immitis
Enterovirus 71

Prion diseases

Streptococcus, group A

Staphylococcus aureus

Group III—Agents with Bioterrorism Potential

■ CDC—Category A

Bacillus anthracis (anthrax)

Clostridium botulinum

Francisella tularensis (tularemia)

Variola major (smallpox) and other pox viruses

Viral hemorrhagic fevers

Arenaviruses

LCM, Junin virus, Machupo virus, Guanarito virus

Lassa Feve

Bunyaviruses

Hantaviruses

Rift Valley Fever

Flaviruses

Dengue

Filoviruses

Ebola

Marburg

Yersinia pestis

List of NIAID Emerging and Re-emerging Diseases 2004

■ CDC—Category B

Brucella species (brucellosis)

Burkholderia pseudomallei (melioidosis)

Burkholderia mallei (glanders)

Coxiella burnetii (Q fever)

Epsilon toxin of *Clostridium perfringens*

Food-borne and Water-borne Pathogens

Bacteria

Campylobacter jejuni

Diarrheagenic E. coli

Listeria monocytogenes

Pathogenic vibrios

Salmonella

Shigella species

Yersinia enterocolitica

Protozoa

Cryptosporidium parvum

Cyclospora cayatanensis

Entamoeba histolytica

Giardia lamblia

Microsporidia

Toxoplasma

Viruses (calciviruses, hepatitis A)

Additional viral encephalitides

California encephalitis

Eastern equine encephalitis

Japanese encephalitis virus

Kyasanur Forest virus

LaCrosse virus

Venezuelan equine encephalitis

Western equine encephalitis

West Nile virus

Ricin toxin (from *Ricinus communis*)

Staphylococcal enterotoxin B

Typhus fever (Rickettsia prowazekii)

■ CDC—Category C

Emerging infectious disease threats such as Nipah virus, additional hantaviruses, and the following pathogens:

Influenza

Other rickettsias

Multidrug-resistant tuberculosis (MDR-TB)

Rabies

Severe acute respiratory syndrome-associated coronavirus

(SARS-CoV)

Tick-borne encephalitis viruses

Tick-borne hemorrhagic fever viruses

Crimean-Congo hemorrhagic fever virus

Yellow fever

Pathogens that naturally emerge with, or are engineered for, increased virulence, increased transmission, and/or the ability to evade the immune response.

Emerging and Re-emerging Infectious Diseases

Severe Acute Respiratory Syndrome— SARS

In the spring of 2003, the world became aware of an outbreak of a newly recognized pneumonia that was named "severe acute respiratory syndrome," or SARS. The outbreak is thought to have begun in southeastern China's Guangdong province in November 2002, with subsequent spread to the special administrative region of Hong Kong by February 2003. Significant outbreaks also occurred in other Asian countries such as Vietnam and Singapore, and in Canada. Epidemiologic investigation showed that the disease disproportionately affected healthcare workers and other close contacts of patients

such as family members. Through an NIAIDsupported contract with Dr. Robert Webster at St. Jude Children's Research Hospital in Memphis, researchers at Hong Kong University and their colleagues at four local hospitals were the first to report to the World Health Organization (WHO) the isolation of a virus that was linked conclusively to SARS patients. Using a highpowered microscope, researchers examined a culture from a lung biopsy sample and found virus particles whose surface was studded with an array of proteins resembling a crown around the virus—a "coronavirus." The researchers then used antibody tests and other molecular tools to confirm that that this deadly coronavirus was present in at least 35 of the SARS patients they were studying.

Before the emergence of SARS (also called SARS-CoV), human coronaviruses were predominately associated with up to 30 percent of common colds. Coronaviruses are the largest single-stranded RNA viruses known and are divided into three serogroups. Recent data indicate that SARS is the prototype strain for a new fourth group of coronaviruses.

In response to the need for rapidly increased research on the SARS coronavirus, in FY 2003, NIAID awarded administrative supplements to grantees to expand activities on the basic biology and immunology of coronaviruses. Currently, NIAID administers more than 20 grants to support basic research on animal coronaviruses and more than 25 grants on SARS. NIAID also supports contracts to develop diagnostics, vaccines, and therapeutics for SARS. In addition, NIAID supports epidemiologic work on SARS and conducts SARS research within its intramural program. Highlights include:

- NIAID Division of Intramural Research (DIR) scientists and their collaborators developed a mouse model of SARS replication that allows the evaluation of vaccines, immunotherapies, and antiviral drugs. Using the model, the scientists demonstrated that mice produce neutralizing antibodies to the SARS virus that protect the mice from re-infection. In addition, transfer of immune sera from infected mice protected other mice from SARS infection. These observations suggest that vaccines that induce neutralizing antibodies and strategies for immunoprophylaxis or immunotherapy are likely to be effective in SARS. Since its development, the mouse model has been used to demonstrate the efficacy of a DNA vaccine, an inactivated vaccine, a vectored vaccine, and three candidate immunotherapies.²⁷
- To date, NIAID screening contracts have evaluated more than 20,000 chemicals for anti-SARS-CoV activity. These NIAIDsupported investigators have screened more

- than 1,400 compounds, including all FDA-approved antiviral drugs. Four compounds, thus far, have shown activity and will be studied further.
- NIAID awarded five contracts, grants, and supplements to companies and researchers towards development of vaccines for SARS, covering a variety of different vaccine approaches.
- NIAID-supported researchers at Hong Kong University have developed a polymerase chain reaction assay for detection of the SARS coronavirus. The test has been shown to detect the SARS virus in respiratory aspirates and fecal samples.
- NIAID is supporting the development of a diagnostic microarray that will be able to detect influenza, SARS, and other respiratory viruses.
- NIAID has expanded its Pandemic Preparedness in Asia contract with St. Jude Children's Research Hospital (Dr. Robert Webster, Principal Investigator) to:
 - Expand efforts to identify the animal reservoirs for coronaviruses in Asia;
 - Establish cell-based laboratory assays to assess the immune response in infected patients; and
 - Conduct seroepidemiologic studies of family members and other close contacts of SARS patients to assess the rates of asymptomatic infections.
- NIAID is supporting the generation and distribution of a variety of SARS reagents for the research community, including:
 - Overlapping peptides. The NIH
 AIDS Research and Reference Reagent
 Program developed important tools
 for researchers to help understand the

immune response to the SARS virus. One of these tools, a set of synthetic overlapping peptides covering the N, M, and S genes of the SARS virus, can be used to map the T cell responses in exposed and infected people. Use of these peptides will help to determine whether SARS patients with different disease outcomes have quantitatively different T cell responses to viral proteins.

- Functional Genomics Resource Center has developed microarrays that contain genetic sequences of SARS coronavirus isolates from the United States, Canada, and Asia, which can be used to detect tiny genetic variations in the SARS viruses. With this information and information on the clinical outcomes of patients infected with SARS, scientists hope to determine which strains are most dangerous and to gain information on the development of antiviral drugs.
- In preparation for the re-emergence of SARS, NIAID is developing a clinical protocol to evaluate Interferon alfacon-1 (INFERGEN), an engineered recombinant interferon molecule that has potent anti-SARS-CoV activity in an *in vitro* assay (cytopathic effect). NIAID's Collaborative Antiviral Study Group is leading these efforts, in consultation with the Centers for Disease Control and Prevention (CDC) and colleagues from Toronto.
- NIAID issued a number of solicitations for grant and contract proposals to further expand research on SARS and accelerate product development. These new programs include development of diagnostics, vaccines, and therapeutics for SARS and will provide opportunities for partnerships between academia, government, and the private sector.

For more information on SARS research updates and opportunities, please visit www.niaid.nih. gov/dmid/sarsopps.htm and www.niaid.nih.gov/factsheets/sars.htm.

West Nile Virus

NIAID supports a robust West Nile virus research portfolio. The following points summarize key research in several different areas:

- NIAID conducts basic research, which leads to a better understanding of the host, pathogen, and environmental factors that influence disease emergence. Basic research determines which flavivirus proteins contribute to the virus's ability to cause disease and examines how protective immune responses are elicited within the central nervous system during acute flavivirus encephalitis.
- A golden hamster model has been developed by NIAID-supported researchers and is used for screening drugs and for examining factors that contribute to immunity. This model has proven useful in evaluating strategies for



Researcher wearing biosafety gear to demonstrate preparation of infected tissue culture plates.

- preventing the complications associated with this emerging infectious disease.
- In 1999, NIAID funded a fast-track project to develop a candidate West Nile virus vaccine with Acambis, Inc. Since then, scientists have developed a prototype vaccine and conducted initial feasibility studies. The vaccine is a chimeric vaccine (West Nile virus protein on a yellow fever vaccine). The Acambis West Nile virus vaccine candidate has so far demonstrated good safety, efficacy, and protection against disease in animal models. Phase I clinical trials in humans began in November 2003 by Acambis.
- A DNA vaccine is being supported by NIAID. A phase I clinical trial is planned for early 2005.
- NIAID has funded investigators to establish
 a system to screen chemical compounds for
 possible antiviral activity against West Nile
 virus. More than 1,000 compounds have been
 screened, and several have moved forward
 to preclinical evaluation. NIAID also is
 supporting research on immunotherapeutics.
- NIAID supports the World Reference Center for Arboviruses, which has reference anti-West Nile sera and seed lots of various strains of West Nile virus. These reagents were provided when requested by investigators in the United States and Canada.
- At the end of FY 2002, the Division of Microbiology and Infectious Diseases (DMID) awarded two contracts to establish Emerging Viral Diseases Centers in response to a Request for Proposals titled "U.S. Based Collaboration in Emerging Viral and Prion Diseases." Each contract establishes broad-based, interactive, multidisciplinary research teams with the scientific expertise needed to study the emergence of a wide variety of zoonotic and arthropod-borne viral pathogens and other emerging viral

- threats. Both contracts also provide capacity to redirect funds and resources in the event of an urgent public health threat from either natural disease or bioterrorist release. Under this provision, research on the newly discovered SARS coronavirus is being conducted at both sites. These contracts cover several important areas of research, including basic biology of the virus, West Nile virus ecology and pathogenic/epidemic potential, diagnosis, prevention, and therapy.
- NIAID supports research aimed at better understanding the vectors of transmission in affected areas. Such an understanding will allow improved monitoring and surveillance for the vectors and the viruses they transmit. NIAID also supports the development and preliminary testing of vector control strategies.
- NIAID intramural scientists also have developed a West Nile virus vaccine candidate, which they have tested in monkeys and horses with promising results. This vaccine candidate is a result of groundbreaking studies conducted more than a decade ago, in which NIAID scientists combined parts of different flaviviruses (a family of viruses that includes West Nile virus, dengue, Japanese encephalitis, and others) to make them weaker and thus more suitable for a live-virus vaccine. The West Nile virus vaccine uses part of the dengue virus as a backbone to which protective antibodyeliciting components of the West Nile virus are added. A supply of this vaccine suitable for human use has been prepared, and phase I human trials are planned for 2005. Development of other vaccine approaches, such as a full-length cDNA-derived West Nile virus vaccine, is under way.

For more information on West Nile virus and NIAID's research portfolio in this area, see www. niaid.nih.gov/publications/wnile/default.htm and www.niaid.nih.gov/factsheets/westnile.htm.

Tuberculosis

M.tb kills more people globally than any other single infectious agent. It is estimated that one-third of the world's population (1.86 billion people) is infected with *M.tb*, and 16.2 million people currently have tuberculosis (TB).²⁸ In 2002, an estimated 8.8 million persons developed TB, and 2 million patients died from this disease. Based on these statistics, TB kills more adults globally than any other single infectious agent.²⁹

The majority of TB cases occur in developing nations. Although TB is essentially a treatable disease, lack of availability of drugs in many countries, poor adherence to treatment schedules due to side effects, and the long duration of treatment (6 to 12 months) have resulted in the development of single drug-resistant and multidrug-resistant TB (MDR-TB) strains producing TB that is much more difficult to cure. Furthermore, the link between HIV and TB is believed to be a major factor in the spread of TB. In 1997, of the 1.86 billion individuals worldwide who were infected with *M.tb*, approximately 10.7 million also were infected with HIV. In Africa, TB cases are increasing by 10 percent each year because of HIV. These factors, combined with a suboptimal public health infrastructure in many countries, contribute to the ongoing spread and re-emergence of TB worldwide.

NIAID TB research efforts are carried out through a comprehensive extramural and intramural program funding all aspects of basic, translational, and applied research leading to a better understanding of TB as an infection and disease, as well as to the development of novel vaccines, drugs, and diagnostics. Research on TB is an important part of NIAID's global health portfolio, as is articulated in the NIAID Global Health Research Plan for HIV/AIDS, Malaria, and Tuberculosis (http://www.niaid.nih.gov/publications/globalhealth/global.pdf). Highlights of the NIAID TB research program include:

 Basic biology. NIAID supports basic research on the disease-causing mechanisms of the bacteria, as well as host immune responses to mycobacterial infection, including the identification of potential targets that might be used in vaccination or treatment of the disease.

- **Drug discovery and evaluation**. NIAID supports the evaluation of new candidate antibiotics against virulent *M.tb* in laboratory assays and in animal models of disease (see http://www.taacf.org). The Institute also supports research to determine mechanisms of drug resistance and activity, early target identification and verification, and clinical testing of prophylactic and therapeutic anti-TB regimens.
- Vaccine development and evaluation.
 NIAID supports basic and clinical research leading to the development and testing of effective new vaccines for the prevention and control of TB.
- Public/private partnerships. NIAID works closely with industry to merge basic science with industry expertise in the development of TB therapeutics and vaccines. NIAID supports these efforts through the award of Challenge Grants, cooperative agreements, and small business grants, as well as participation in public-private partnerships.

Lyme Disease

Lyme disease (borreliosis) is the most prevalent tick-borne infectious disease in the United States. In 2003, the CDC reported 21,273 cases throughout the United States.³⁰ Five states—New York, Pennsylvania, Massachusetts, Connecticut, and New Jersey—accounted for 80 percent of all reported cases.³¹

The major goals of the NIAID Lyme disease research program are to develop better means of diagnosing, treating, and preventing this disease. To accomplish these objectives, the NIAID Lyme disease research portfolio includes a broad range of activities that are essential to

increasing our understanding of the disease. The studies include both intramural and extramural research on animal models of disease, microbial physiology, molecular and cellular mechanisms of pathogenesis, mechanisms of protective immunity, vectors and disease transmission, efficacy of different modes of antibiotic therapy, and development of more sensitive and reliable diagnostic tests for both early (acute) and late (chronic) Lyme disease.

NIAID intramural investigators are studying Lyme disease on the NIH campus in Bethesda, Maryland, and at the Rocky Mountain Laboratories (RML) in Hamilton, Montana, where NIAID scientists discovered the etiologic agent Borrelia burgdorferi in the early 1980s.³² RML scientists are using microarray technology to identify genes associated with unique aspects of the pathogenicity of Lyme disease and other relapsing fever microorganisms. In addition, working with members of the California Department of Health Services, RML researchers identified two new areas in California where a tick-borne relative of the Lyme disease bacterium called Borrelia hermisii has been found to cause relapsing fever. This finding will help alert the medical community and public health workers about the potential for this tick-borne disease to be acquired in these areas.

On the NIH campus, NIAID clinical investigators seek to better understand the natural history of chronic Lyme disease and possible causes for persisting symptoms. To this end, two clinical studies currently are ongoing at the NIH Clinical Center: one to evaluate and treat patients with classic Lyme disease, and the other to conduct a comprehensive clinical, microbiologic, and immunologic assessment of patients who have suspected chronic Lyme disease despite previous antibiotic therapy. In addition, NIAID clinical scientists are collaborating with colleagues to further evaluate C6 peptide enzyme-linked immunosorbent assay (ELISA), which can detect infection with both the U.S. and European strains of Borrelia, and can be used to diagnose Lyme

disease in patients who have received the Lyme disease vaccine.

NIAID also supports an FY 2003 research initiative titled Partnerships for Hepatitis B and Vector-Borne Diseases. This initiative is an expansion of previous efforts that targeted animal vectors of disease and fostered development of partnerships among government, academia, and the biotechnology and pharmaceutical industries. NIAID funded two grants on Lyme disease immunology and vaccine development under the 2003 initiative.

Lyme borreliosis and ehrlichiosis will continue to be areas of high priority for basic research for NIAID, especially with regard to: (1) the characterization and treatment of acute and chronic infection; (2) the influence of co-infection with other vector-borne pathogens on the diagnosis, treatment, and severity of Lyme disease; and (3) the development of rapid, sensitive, and specific diagnostic tests and preventive strategies (e.g., vaccines and vector control measures).

Influenza

In the United States, pneumonia and influenza are the seventh leading cause of death, responsible for 2.7 percent of all deaths.³³ Research supported by NIAID has led to many new insights about how influenza causes disease.

The major goal of the NIAID influenza program is to support research leading to more effective approaches to control influenza virus infections. NIAID currently supports research in the following major areas:

- Basic biology. NIAID supports basic research on virus structure and function, viral pathogenesis, and the host response to infection.
- Surveillance/epidemiology. NIAID supports research to better understand the natural history and emergence of influenza

- viruses with pandemic potential and to evaluate community-based strategies for interrupting the spread of influenza.
- Public/private partnerships. In 2000, NIAID awarded three challenge grants (requiring matching funds) to private-sector companies for the development of new vaccines against pandemic influenza strains. These companies are using live-attenuated viruses, virus-like particles, tissue culture substrates, and reverse genetics strategies to rapidly produce high-growth viruses for vaccine production. In 2003 and 2004, NIAID awarded multiple cooperative agreements to private sector companies/ academic collaborators to develop new countermeasures against influenza virus.
- **Drug discovery and evaluation.** NIAID supports the development of novel drugs against influenza and the evaluation of these new agents in both *in vitro* screening assays and animal models. In 2004, NIAID screened more than 170 compounds.
- Vaccine development and evaluation.

 Developing new influenza vaccines and strategies has been a major focus of the NIAID influenza program. These strategies include supporting the development of live-attenuated and recombinant vaccines, immunomodulators and adjuvants, cell culture-based vaccines, and basic research aimed at optimizing the immune response.

 NIAID also supports the production of pilot lot vaccines against avian influenza subtypes of high pandemic potential.

In 2003, FDA approved FluMist, a new intranasally administered influenza vaccine. NIAID supported the development of this vaccine for more than 30 years, through both its intramural program and extramural contracts and cooperative research and development agreements (CRADAs). Since the 1970s, NIAID has supported studies to evaluate the safety and

immunogenicity of strains of the influenza virus that replicate only in the upper respiratory tract (temperature sensitive) and cause only minimal symptoms (attenuated), which could therefore be used as vaccines. In 1995, NIAID entered into a CRADA with Aviron for the commercialization of the product. Over the past 8 years, NIAID has conducted a series of clinical trials to evaluate the vaccine in children and in HIV-positive adults and children. In June 2003, FDA approved FluMist for healthy children and adults aged 5 to 49 years. The potential advantages of FluMist include the ease of administration and the ability to induce a broad immune response.

In addition, NIAID's intramural program is developing vaccines against potential pandemic influenza strains. An optimal public health response in the event of a potential pandemic requires that vaccines be available with minimum delay. NIAID intramural scientists and colleagues from CDC have initiated a collaborative, proactive approach to pandemic preparedness. They plan to generate and evaluate up to two dozen candidate vaccines against influenza A subtypes that are recognized to have pandemic potential. Using classic genetic recombination techniques, the team has developed a candidate inactivated H9N2 vaccine that has demonstrated efficacy in preclinical testing. Clinical studies of this candidate vaccine are planned for 2005. In addition, the scientists have engineered liveattenuated vaccine candidates against the 1997, 2003, and 2004 H5N1 avian viruses. In 2004, the H5N1 virus killed 31 of 43 humans known to be infected as of October 4th.34

In 2003, NIAID also expanded its Pandemic Preparedness in Asia contract with St. Jude Children's Research Hospital in Memphis. (The original award was made in 1998 for the surveillance and characterization of avian influenza viruses with pandemic potential in the live bird markets in Hong Kong.) Activities conducted under this expansion include establishing animal influenza surveillance sites in Asia, generating high-yield vaccine candidates

against influenza strains with pandemic potential and accompanying reagents, supporting an international animal surveillance training course in Hong Kong (March 2004), and studying newly emerging influenza strains infecting swine in the United States.

The major focus of the NIAID influenza program will continue to be on basic and applied research that promises to further the development of new and improved vaccines and antiviral agents.

For more information on influenza, including weekly reports on flu activity, go to www.cdc.gov/flu/weekly/fluactivity.htm.

Prion Diseases

NIAID's DIR has a productive and growing program focused on transmissible spongiform encephalopathies (TSEs). These diseases also are called "prion diseases" because they are believed to be caused and transmitted by prion proteins, a new type of infectious agent discovered in the 1980s. Prion proteins enter cells and cause normal cellular proteins to adopt abnormal threedimensional structures, which in turn leads to disease. TSEs are fatal neurodegenerative diseases and include scrapie, Creutzfeldt-Jakob disease (CJD), bovine spongiform encephalopathy (BSE or "mad cow" disease), and chronic wasting disease (CWD) of deer and elk. Since the onset of the BSE epidemic in the United Kingdom in the 1980s, the disease has resulted in the destruction of millions of animals in Europe. Because the BSE epidemic was temporally and geographically associated with the emergence of a variant form of CJD in humans, health officials believe the disease was spread to humans by infected beef. In May 2003, the finding of BSE in a single cow in Canada resulted in a ban on exportation of certain live ruminants and ruminant products from Canada to the United States.

NIAID's intramural TSE research is aimed at increasing fundamental understandings of prion protein (PrP) and the mechanisms responsible for

the accumulation in nervous system cells of the abnormal form of PrP, which appears to underlie TSE pathogenesis. Studies also are ongoing to elucidate the mechanisms of cross-species transmission of TSE disease. This work is highly important in light of the epidemiology of variant CID as well as the discovery of CWD in deer and elk herds beyond the areas in the western United States in which it was long known to exist. DIR scientists have conducted experiments that suggest that species once thought to be resistant to certain TSE strains can serve as lifelong carriers of the infection without ever becoming sick. Infrastructure improvements at NIAID's RML, including construction of new BSL-3 laboratory and animal facilities, have allowed expansion of TSE transmission studies. An important study to determine whether CWD PrP can be transmitted to nonhuman primates via oral or intracerebral routes has begun, using CWD samples obtained through NIAID's collaboration with the Wyoming Department of Health.

DIR scientists also have developed a high-throughput screening method to find compounds that show promise as potential TSE therapeutics. Studies of the potential use of antibody and other vaccine-based therapies for TSEs are ongoing in NIAID laboratories. In addition, to advance earlier peptide studies, novel PrP peptides have been synthesized and are being evaluated for their ability to block the conversion of normal PrP to abnormal PrP *in vitro*. PrP peptides dispensed by direct injection or delivered by gene therapy might provide specific therapeutic treatment for TSE diseases. Promising compounds will be evaluated *in vivo* through a DMID contract with Utah State University. (See below.)

NIH provides grant support for investigatorinitiated studies of CWD transmission that seek to better understand prion entry, trafficking, and neuroinvasion in the lymphoid system, which could provide a basis for development of diagnostic and intervention strategies. In addition, NIH has taken a number of actions in response to the Department's 2001 BSE/TSE Action Plan, including:

- NIAID awarded a 7-year, \$8.4 million contract to Colorado State University to establish an emerging disease research center focused on CWD. Researchers at the center will isolate, identify, and characterize strains of CWD; evaluate the potential for both inter- and intraspecies transmission; study the pathogenesis of CWD; perform preclinical, animal model-based evaluation of newly developed prevention measures or therapies for CWD; develop, analyze, and distribute reagents, infectious material, molecular clones, and transgenic mice to the research
- community; and implement a systematic approach to furthering understandings of the ecologic and environmental factors influencing the emergence, spread, and distribution of CWD and of the basic epidemiology and clinical aspects of these diseases.
- NIH is evaluating potential anti-TSE compounds in animal models. Through expansion of an NIAID contract with Utah State University, candidate compounds are evaluated for efficacy in transgenic animals that have a shortened time to death. This model was established at Utah State in collaboration with NIAID's RML.